Hauda, K.

1632

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RAW SEQUENCE LISTING PATENT APPLICATION US/08/982,284

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This Raw Listing contains the General Information Section and up to the first 5 pages.

1 2		SEQUENCE LISTING		
3	(1) G	eneral Information: ENTE		
4	(_,	LIVI		
5	(i)	APPLICANT: LUBON, Henryk		
6		PALEYANDA, Rekha		
7		DROHAN, William		
8		VELANDER, William		
9				
10	(11)	TITLE OF INVENTION: METHODS FOR THE DEGRADATION AND	DIE	
11		DETOXIFICATION OF ORGANIC MATERIAL USING URINE PRODUCED	BY	
12 13		TRANSGENIC ANIMALS AND RELATED TRANSGENIC ANIMALS AND PROTEINS		
14		PROTEINS		
15	(iii)	NUMBER OF SEQUENCES: 3	00 APR	ECH 12
16	(111)	norman or bigoincis. 3	2	
17	(iv)	CORRESPONDENCE ADDRESS:	20	
18	,	(A) ADDRESSEE: FOLEY & LARDNER	9-	-417
19		(B) STREET: 3000 K Street, N.W.	S	- Divi
20		(C) CITY: Washington		-4-5
21		(D) STATE: D.C.	<u> </u>	30
22		(E) COUNTRY: U.S.A.	က်	:
23		(F) ZIP: 20007-5109	9: 35	1/2
24			S	1500/2900
25	(v)	COMPUTER READABLE FORM:		9
26		(A) MEDIUM TYPE: Floppy disk		
27		(B) COMPUTER: IBM PC compatible		
28 29		(C) OPERATING SYSTEM: PC-DOS/MS-DOS		
29 30		(D) SOFTWARE: PatentIn Release #1.0, Version #1.30		
31	(37i)	CURRENT APPLICATION DATA:		
32	(VI)	(A) APPLICATION NUMBER: US 08/982,284		
33		(B) FILING DATE: 01-DEC-1997		
34		(C) CLASSIFICATION:		
35		(-,		
36	(viii)	ATTORNEY/AGENT INFORMATION:		
37		(A) NAME: Granados, Patricia D.		
38		(B) REGISTRATION NUMBER: 33,683		
39		(C) REFERENCE/DOCKET NUMBER: 030523/0141		
40				
41	(ix)	TELECOMMUNICATION INFORMATION:		
42		(A) TELEPHONE: (202) 672-5300		
43		(B) TELEFAX: (202) 672-5399		
44				
45 46	(2) TMEO	PMATTON FOR CEO ID NO.1.		
46	(Z) INFO	RMATION FOR SEQ ID NO:1:		

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RAW SEQUENCE LISTING PATENT APPLICATION US/08/982,284

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47		
48	(i) SEQUENCE CHARACTERISTICS:	
49	(A) LENGTH: 69 base pairs	
50	(B) TYPE: nucleic acid	
51	(C) STRANDEDNESS: single	
52	(D) TOPOLOGY: linear	
53	(2) 10101001	
54	(ii) MOLECULE TYPE: DNA (genomic)	
55	(II) Modecold IIIE. DNA (genomic)	
56		
57		
58	(wi) CHOMPINGE PERGETERION, CHO. ID. NO. 1.	
59	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:	
60		
61	GGTACGTCGA CACAGGTGCC AGTGCCTCCA GAATGTGGCA GCTCACAAGC CTCCTGCTGT	60
62		
63	TCGTGGCCA	69
64		
65	(2) INFORMATION FOR SEQ ID NO:2:	
66	•	
67	(i) SEQUENCE CHARACTERISTICS:	
68	(A) LENGTH: 22 base pairs	* · 1
69	(B) TYPE: nucleic acid	00
70	(C) STRANDEDNESS: single	OO APR-6
71	(D) TOPOLOGY: linear	哥哥
72		为一部
73	(ii) MOLECULE TYPE: other nucleic acid	
74	(A) DESCRIPTION: /desc = "Primer"	S 200
75	(4, -4, -4, -4, -4, -4, -4, -4, -4, -4, -	Eng. 635
76		
77		≥ بي
78	·	(a) 5
79	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:	M 9: 35
80	(AI) Digolaci Discalliton. Dig ID No.1.	ç
81	CCCAGGCTCA GCGCACTCAT CC	22
82	CCASGCTCA GCGCACTCAT CC	
83	(2) INFORMATION FOR SEQ ID NO:3:	
84	(2) INFORMATION FOR SEQ 1D NO.3.	
85	(i) SEQUENCE CHARACTERISTICS:	
	(A) LENGTH: 21 base pairs	
86	(B) TYPE: nucleic acid	
87		
88	(C) STRANDEDNESS: single	
89	(D) TOPOLOGY: linear	
90		
91	(ii) MOLECULE TYPE: other nucleic acid	
92	(A) DESCRIPTION: /desc = "Primer"	
93		
94		
95		
96		
97	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:	
98		
99	GTCACAGCAA TGCCACCTGA C	21

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SEQUENCE VERIFICATION REPORT PATENT APPLICATION US/08/982,284

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